

FIG. 2

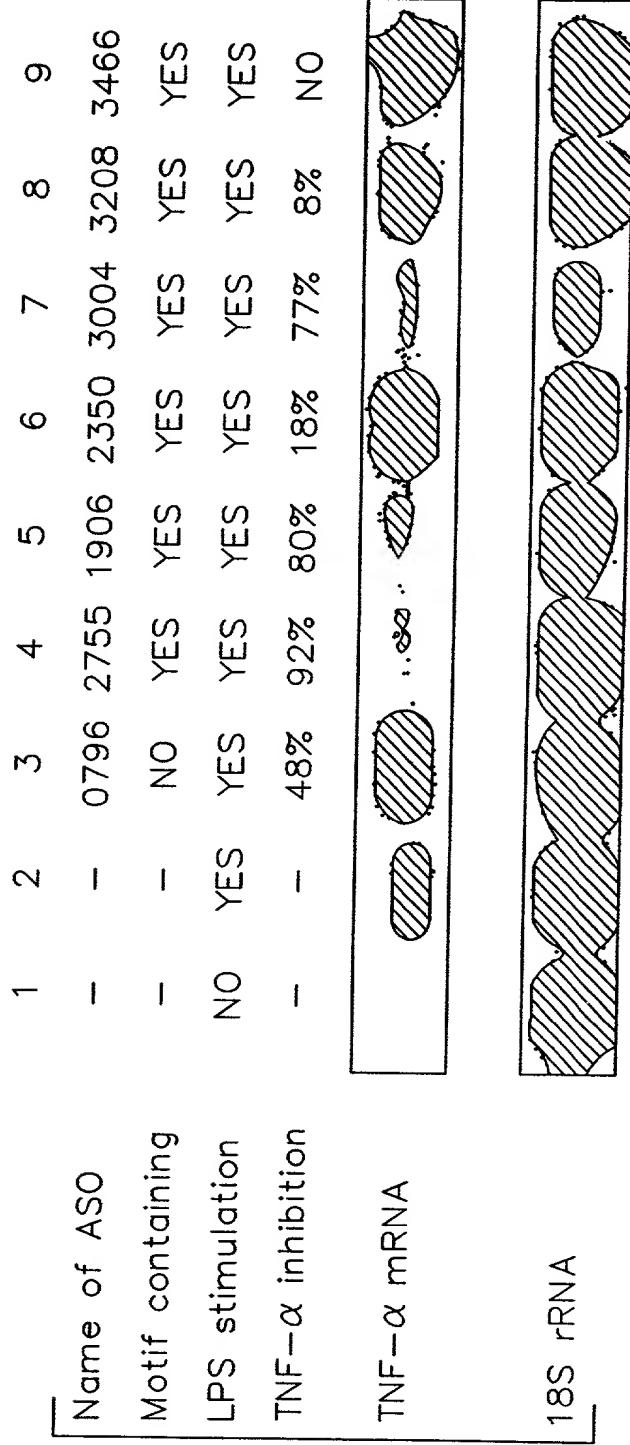


Fig. 3

1 gaattccggg t~~gat~~ttcact cccggctgtc caggcttgc caccaggccc acccaggcc
61 tcctgaggg tcaaggccgtc cacaaggccc ccagctcc ttcttcgac gacccaaaca
121 caggcctcg gactcaaac agctttccc tccaaccgg tttctctcc tcaacggact
181 cagcttcg aaggccctcg cagttcttagt tctatcttt tcctgcattc tgtctggaaag
241 tt~~g~~aaaaggaa acagaccaca gacctggcc caaaaagaaa cggggcaat tgaggcaat aggttttgaaag
301 gg~~g~~catggg AcggatGGG g~~c~~ctccagg~~g~~ tcctacacac aaatcagtca g~~t~~ggcccaga
361 agacccctcg cg~~a~~atcgga g~~c~~GGAGtg~~g~~ g~~g~~gggtatc ct~~t~~gatgtct
421 gtgttcccc aacttccaa atccccggcc cggcgatgg~~a~~ g~~a~~ggaaaccc g~~t~~ggcccaga
481 tg~~c~~aggccc actaccgctt cttccagatg agctcatgg~~t~~ tttctccacc aaggaaagt~~t~~
541 tc~~c~~gctgggtt gaatgattt tccccggcc tccctcgcc ccAGGAcat ataaaggcag
601 tt~~g~~ttggcac acccaggccg cagacgctcc ctcagcaagg acaggagg~~g~~ accaggtaag
661 aGGGA~~g~~agaa g~~c~~aaactacag accccccctg aaaacaacc tcagacgca catccccctga
721 caaggctgc~~a~~ ggcagg~~t~~tt cttcctctca catactgacc cacggcttca ccctctcc
781 cctggaaagg acaccatgag cactgaaagg atqatccGGG Acgttggagct ggcggaggag
841 gg~~c~~tccca~~a~~ gaaagacagg agggccccag ggc~~t~~ccaggc ggtgcttgc cctcaggcc
901 ttctccccc t~~g~~atcgtgg~~c~~ acggccacc acqctctt gcctgctga ctttggaaqt~~g~~
961 atcgcccccc agaGGGA~~g~~aa ggtgagtgcc tg~~g~~ccaggct tcatccactc tcccacccaa
1021 gg~~G~~GAat~~g~~a gagacggcaag agaGGGA~~g~~ag~~g~~ agatGGGA~~g~~at~~g~~ ggtgaaagat~~g~~
1081 aGGGAGGGAt g~~g~~ag~~g~~aaa~~g~~aaa aaaaacatgg~~a~~ gaaaggacgg~~g~~ GAtggcagaaa
1141 aaggagatGGG GA~~a~~gag~~g~~ag~~g~~ ag~~g~~aaaagat~~g~~ g~~g~~ag~~g~~acag~~g~~ gatgtctggc
1201 tg~~c~~tactaa~~a~~ g~~t~~gtgtatgg agtgaat~~g~~aa tgaat~~g~~aaat~~g~~ aatgaacaag~~g~~ acatggaaagg
1261 aataagat~~a~~at gg~~g~~agacagat g~~t~~gggggtgt~~g~~ ag~~g~~aaaggag~~g~~aa tggGGGA~~g~~aa~~g~~ acaaaqt~~g~~at~~g~~
1321 at~~g~~aaataa~~g~~ ag~~g~~gtgagac GGA~~a~~aat~~g~~at~~g~~aa cayctaagg~~a~~ g~~g~~ag~~g~~atgg~~g~~
1381 gg~~g~~gataa~~g~~ ag~~g~~aaaggaa~~g~~ ag~~g~~aaaggaa~~g~~ tggcacac~~g~~ aq~~g~~acactca~~g~~ GGAAaggagg
1441 tg~~t~~gaaat~~g~~ g~~c~~taaggag~~g~~ t~~g~~g~~g~~aaagg~~g~~at~~g~~ g~~g~~atgg~~g~~ag~~g~~ agaaaacc~~g~~ acaccc~~t~~cagg
1501 gcttaaccat tctcc~~t~~tc~~c~~ caggccag~~c~~ c~~g~~ggccag~~c~~ ag~~g~~ggccag~~c~~ ct~~g~~ttcc~~t~~cc~~c~~ t~~t~~taagg~~g~~gt~~g~~ actccctc~~g~~ga~~c~~ t~~c~~t~~c~~t~~c~~aa~~t~~ tca~~g~~cc~~c~~c~~t~~ct
1561 t~~g~~ttaaccat tctcc~~t~~tc~~c~~ cccaa~~c~~agg~~t~~ cccaa~~c~~agg~~t~~ cccaa~~c~~agg~~t~~ cccaa~~c~~agg~~t~~ cccaa~~c~~agg~~t~~

Fig. 4A

Fig. 4B

3241 tttaaaaata ttatatctgat taagttgtct aaacaatgct gatttggta ccaaactgtca
3301 ctcatgtcg agccctctgct ccccaggGGA gttgtgtctg taatcgccct actattcagt
3361 ggccgagaaat aaagtttgtct tagaaaaagaa acatggtctc cttttggaa ttaattctgc
3421 atctgcctct tc ttgtgggt tc ccctaagt GGAAGaaAGC tc tctccac aggtttaag
3481 atccctcgga ccagtc ca tc tttagact cctaggccc tgagacct acataaaca
3541 agcccaacag aatattccc atccccagg aaacaagg ctgaaaggc ttacacctcc
3601 ctcaaggcat GGAAattttcc aactctGGGA atcc

Fig. 4C

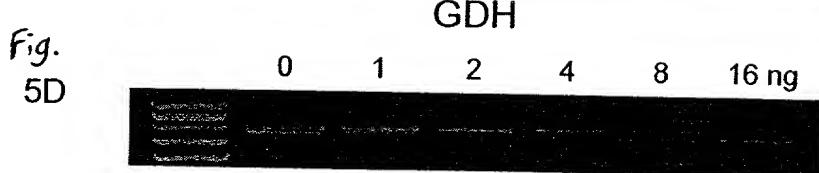
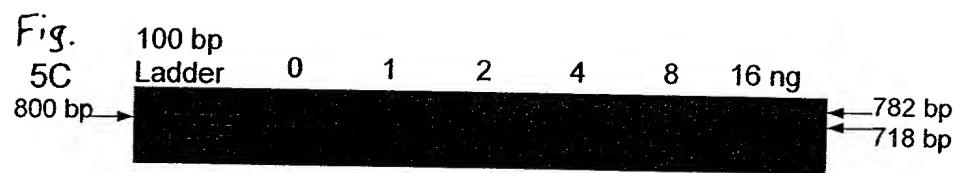
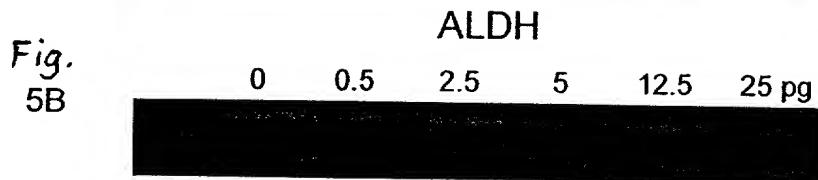
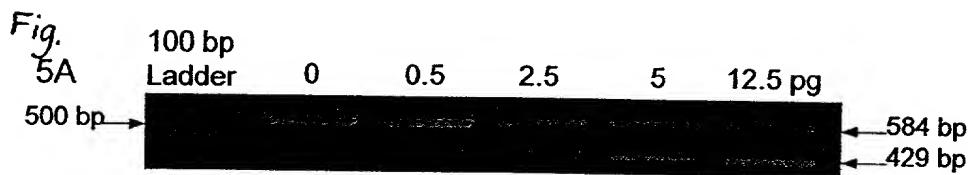
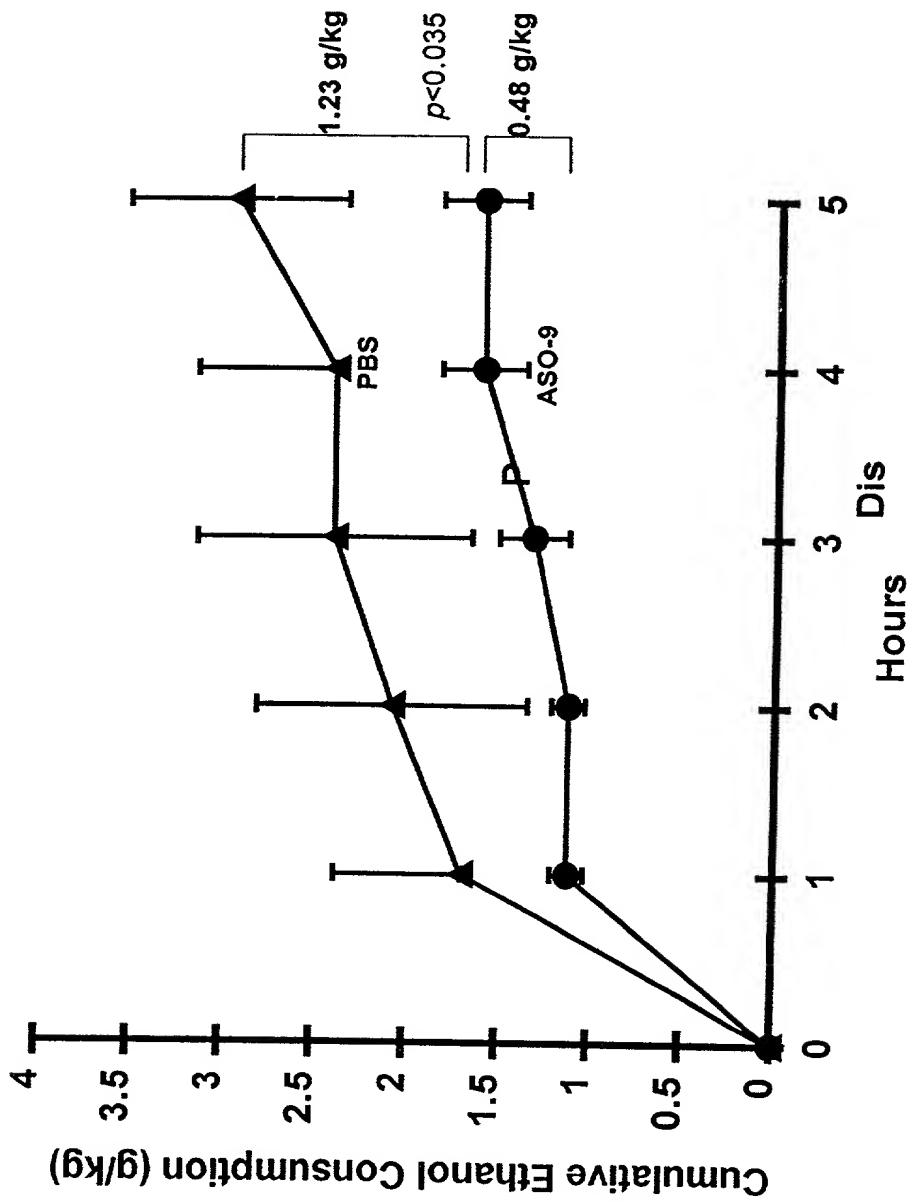


FIGURE 6



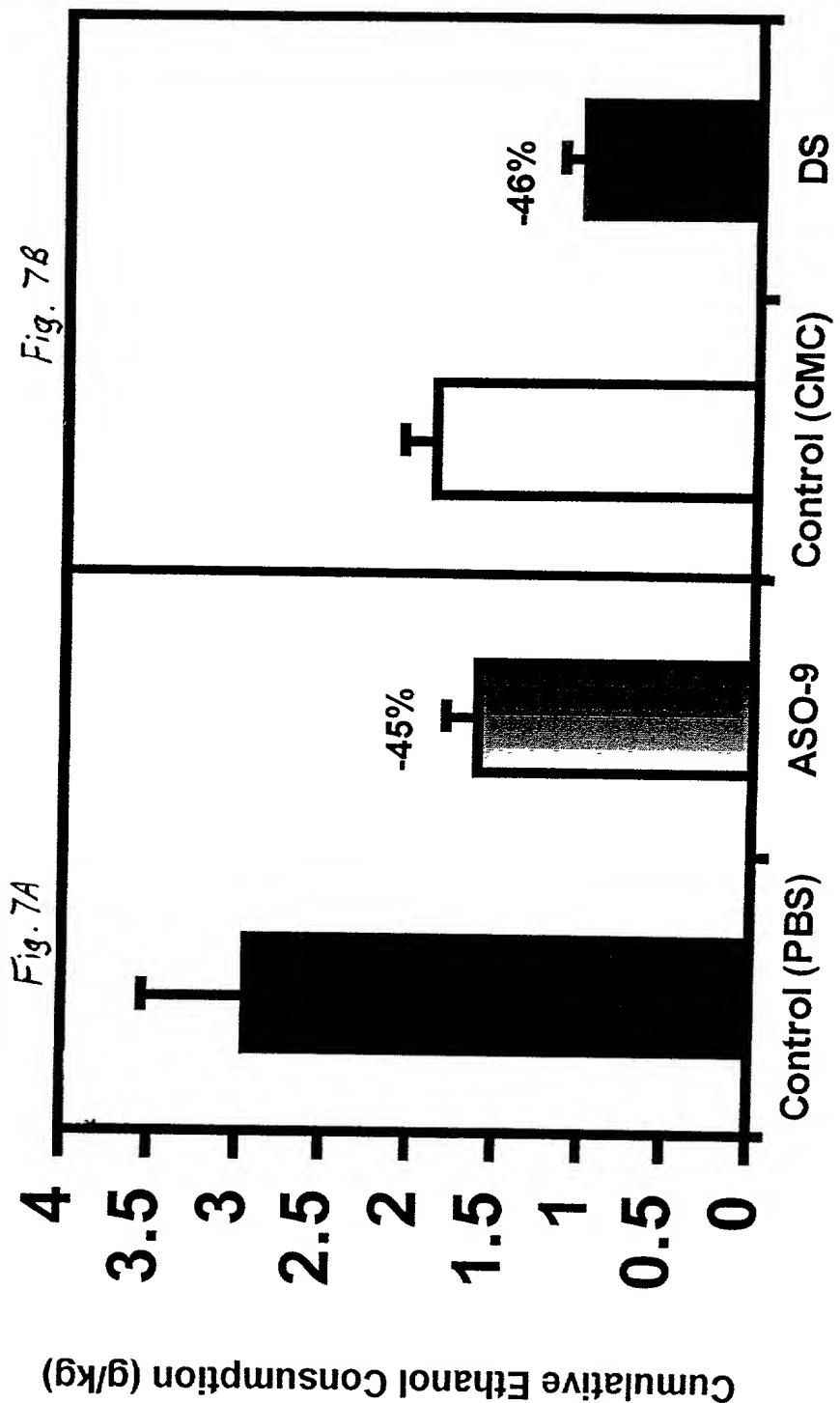
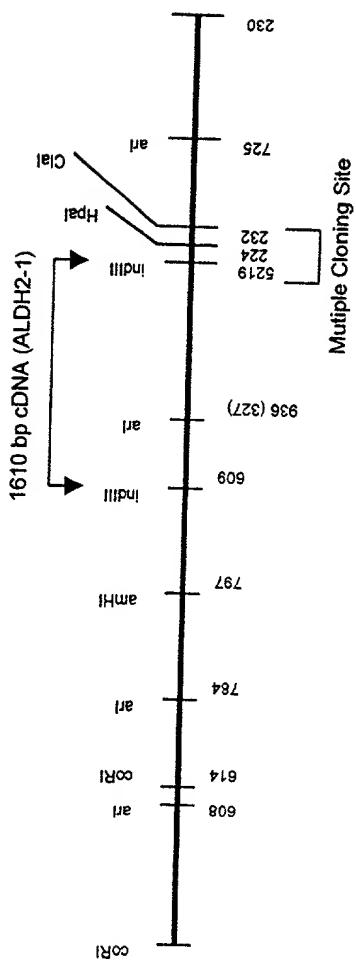


Fig. 8A



The diagram illustrates the 1610 bp cDNA (ALDH2-2) with various restriction enzyme cleavage sites indicated by arrows. The cDNA is represented as a horizontal line with two arrows pointing from left to right, indicating the direction of transcription. Key features include:

- Top Panel:** A bracket labeled "+ 400 bp 3' UTR" spans the entire length of the cDNA. Within this region, several restriction sites are marked: **Cla**I at position 6284, **Hpa**I at 5801, **Hind**III at 5778, and **Nar**I at 5793.
- Bottom Panel:** A bracket labeled "1610 bp cDNA (ALDH2-2)" covers the first 1610 bp. Within this region, restriction sites are marked: **Eco**RI at 1608, **Nar**I at 1614, **Eco**RI at 1897, and **Hind**III at 3778. The distance between the **Hind**III site at 3778 and the **Nar**I site at 5793 is labeled as 4105 (327).
- Right Side:** A vertical column labeled "Multiple Cloning Site" lists additional restriction sites: **Eco**RI at 1, **Nar**I at 6284, and **Cla**I at 6284.

Fig. 8B

FIGURE 9

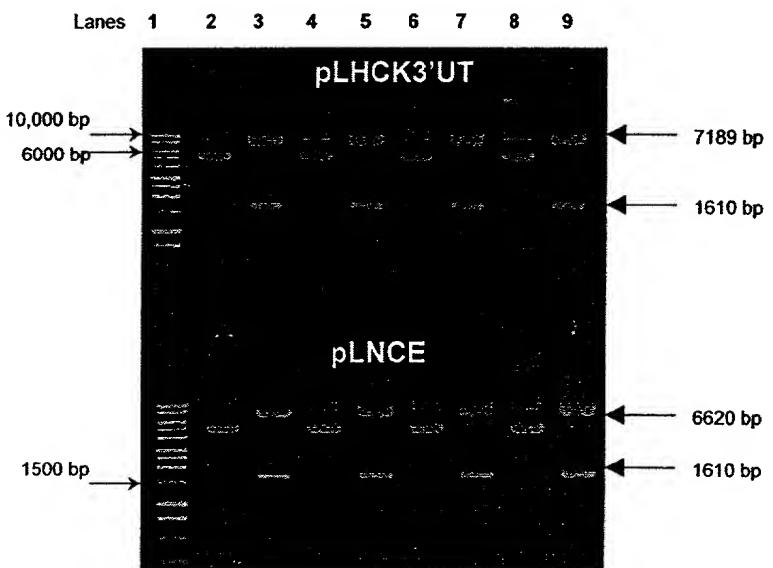
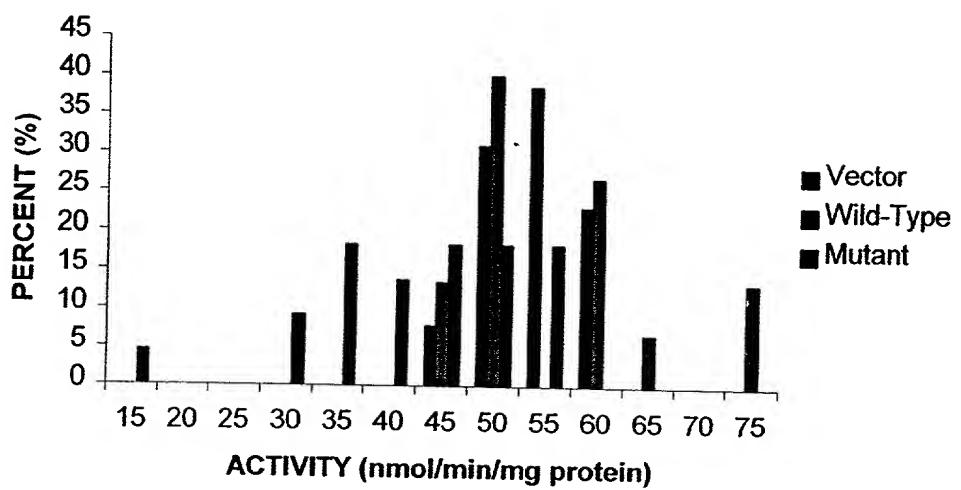


FIGURE 10

H4-II-E-C3 TRANSDUCTION



GCTTTATCTG CTAAGCTCCG CTCAGTTTAG CATGCTGCAG
GCCGCACTCA GCACCGCCCC CGGTGGGCCA CGCCTGAGCC
GCCTGCTGTC CGCCGCCGCC ACCAGCGCGG TGCCAGCCCC
CAACCAGCAG CCCGAGGTCT TCTGCAACCA GATCTTCATT
AACAAATGAGT GGCATGATGC TGTAGCAAG AAAACATTCC
CCACCGTCAA CCCTTCCACG GGGGAGGTCA TCTGCCAGGT
AGCCGAAGGG AACAAGGAGG ACGTAGACAA GGCAGTGAAG
GCCGCTCAGG CAGCCTTCCA GCTGGGCTCG CCCTGGCGCC
GCATGGATGC ATCTGACAGG GCCGGGCTGT TGTACCGATT
GGCTGATCTC ATCGAACGGG ACCGGACCTA CCTGGCGGCC
TTGGAGACCC TGGACAACGG CAAGCCTTAT GTCATCTCCT
ACCTGGTGA TTTGGACATG GTTCTGAAAT GTCTCCGCTA
TTATGCTGGC TGGGCTGACA AGTACCAACGG GAAAACCATT
CCCATCGATG GCGACTTCTT CAGCTACACC CGCCACGAGC
CTGTGGGCGT GTGTGGACAG ATCATTCCGT GGAACCTCCC
GCTCCTGATG CAAGCCTGGA AGCTGGGCC TGCCCTGGCA
ACTGGAAACG TGGTGGTGAT GAAAGTGGCC GAGCAGACAC
CGCTCACTGC ACTCTACGTG GCCAACTTGA TCAAGGAGGC
AGGCTTCCCC CCTGGTGTGG TCAATATTGT TCCTGGATT
GGCCCTACCG CCGGGGCTGC CATCGCGTCC CACGAGGATG
TGGACAAAGT GGCTTCACA GGTTCCACTG AGGTTGGTCA
CCTAATCCAG GTGCCGCCG GGAGCAGCAA TCTCAAGAGA
GTAACCTGG AACTGGGGGG AAAGAGCCCC AATATCATCA
TGTCAAGACGC TGACATGGAC TGGGCTGTGG AACAGGCCA
CTTGCCCTG TTCTTCAACC AGGGCCAGTG CTGTTGTGCG
GGCTCCCGA CCTCGTGCA GGAGGATGTG TATGATGAAT
TCGTGGAACG CAGTGTGGCC CGGGCCAAGT CTCGGGTGGT
CGGGAACCT TTGACAGGCC GGACGGAGCA GGGGCCGAG
GTGGATGAGA CTCAGTTAA GAAGATCCTG GGCTATATCA
AGTCAGGACA ACAAGAAGGG GCGAAGCTGC TGTGCGGTGG
GGGCGCCGCC GCAGACCGTG GTTACTTCAT CCAGCCCACC
GTGTTGGAG ACGTCAAAGA TGGCATGACC ATCGCCAAGG
AGGAGATCTT CGGACCAGTG ATGCAGATCC TCAAATTCAA
GACCATTGAG GAGGTTGTGG GCGGAGCCAA TAATTCCAAG
TACGGGCTGG CTGCCGCTGT CTTCACAAAG GACCTGGACA
AGGCCAATTAA CCTGTCCTAA GCTCTGCAGG CTGGGACTGT
GTGGATCAAC TGCTACGATG TGTGTTGGGC CCAGTCCCCA
TTTGGTGGCT ATAAGATGTC GGGGAGCGGC AGGGAGCTGG
GCGAGTATGG CCTGCAGGCC TACACGGAAG TGAAGACGGT
CACCGTCAA GTGCCACAGA AGAACTCGTA AAGTGGCGTG

Fig. 11A

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CAGGCTTCCT CAGCCAGCGC CCAAAAACCC AACAAAGATCC
TGAGAAAAGC CACCACCAAG CACACTGCAG CTGCCAAGAG
AAAACCCCTT CACCAAAGCG TCTTGGGCCA AGAAAGTCAG
GATTTGATAA ACAGGGCAGG GTTGGTGGGC GGTGTGTGGG
GAGCATCCCA GTAAAATGGG GAAGGGAGGA GCTCTGTGCA
GACTACCACG CGCACGCACA CACGCTCACT GGGCCTTCT
GTGCTGGATG CTGGTTCCAC CCTCAGTGCT TAAACAAATG
AGCAATAAA

Fig. 11B

GCTCTCGGTC CGCTCGCTGT CCGCTAGCCC GCTGCGATGT
TGC CGC CTGC CGCC GCGCTCGG GCCCGCCTG GCCGCCGCCT
CTT GTCA GAGCC GCCGCCACCC AGGCCGTGCC TGCCCCAAC
CAGCAGCCCCG AGGTCTTCTG CAACCAGATT TTCATAAACAA
ATGAATGGCA CGATGCCGTC AGCAGGAAAA CATCCCCAC
CGTCAATCCG TCCACTGGAG AGGTCATCTG TCAGGTAGCT
GAAGGGACA AGGAAGATGT GGACAAGGCA CGTGAAGGCC
GCCCGGGCGC CTCCAGCTG GGCTCACCTT GGCGCCGCAT
GGACGCATCA CACAGCGGCC GGCTGCTGAA CC CGCTGGCC
GATCTGATCG AGCGGGACCG GACCTACCTG GC GGCC TTGG
AGACCCTGGA CAATGGCAAG CCCTATGTCA TCT CCTACCT
GGTGGATTG GACATGGTCC TCAAATGTCT CCGGTATTAT
GCCGGCTGGG CTGATAAGTA CCACGGGAAA ACCATCCCCA
TTGACGGAGA CTTCTTCAGC TACACACGCC ATGAACCTGT
GGGGGTGTGC GGGCAGATCA TTCCGTGGAA TTTCCCGCTC
CTGATGCAAG CATGGAAGCT GGGCCCAGCC TTGGCAACTG
GAAACGTGGT TGTGATGAAG GTAGCTGAGC AGACACCCCT
CACCGCCCTC TATGTGGCCA ACCTGATCAA GGAGGCTGGC
TTTCCCCCTG GTGTGGTCAA CATTGTGCCT GGATTGGCC
CCACGGCTGG GCCCGCCATT GCCTCCCATG AGGATGTGGA
CAAAGTGGCA TTCACAGGCT CCACTGAGAT TGGCCGCGTA
ATCCAGGTTG CTGCTGGGAG CAGCAACCTC AAGAGAGTGA
CCTTGGAGCT GGGGGGAAG AGCCCCAAC A TCATCATGTC
AGATGCCGAT ATGGATTGGG CCGTGGAAACA GGCCCACCTC
GCCCTGTTCT TCAACCAGGG CCAGTGCTGC TGTGCCGGCT
CCCGGACCTT CGTGCAGGAG GACATCTATG ATGAGTTGT
GGTGC GGAGC G TTGCCC GGG CCAAGTCTCG GGTGGTCGGG
AACCCCTTG ATAGCAAGAC CGAGCAGGGG CCGCAGGTGG
ATGAAACTCA GTTTAAGAAG ATCCTCGGCT ACATCAACAC
GGGGAAGCAA GAGGGGGCGA AGCTGCTGT TGGTGGGGC
ATTGCTGCTG ACCGTGGTTA CTTCATCCAG CCCACTGTGT
TTGGAGATGT GCAGGATGGC ATGACCATCG CCAAGGAGGA
GATCTCGGG CCAGTGATGC AGATCCTGAA GTTCAAGACC
ATAGAGGAGG TTGTTGGGAG AGCCAACAAT TCCACGTACG
GGCTGGCCGC AGCTGCTTC ACAAAGGATT TGGACAAGGC
CAATTACCTG TCCCAGGCC TCCAGGCGGG CACTGTGTGG
GTCAACTGCT ATGATGTGTT TGGAGCCCAG TCACCCTTG
GTGGCTACAA GATGTGGGG AGTGGCCGGG AGTTGGCGA
GTACGGGCTG CAGGCATACA CTGAAGTGA AACTGTCACA
GTCAAAGTGC CTCAGAAGAA CTCATAAGAA TCATGCAAGC

Fig. 12A

TTCCTCCCTC	AGCCATTGAT	GGAAAGTTCA	GCAAGATCAG
CAACAAAACC	AAGAAAAATG	ATCCTTGCCT	GCTGAATATC
TGAAAAGAGA	AATTTTCCT	ACAAAATCTC	TTGGGTCAAG
AAAGTTCTAG	AATTTGAATT	GATAAACATG	GTGGGTTGGC
TGAGGGTAAG	AGTATATGAG	GAACCTTTA	AACGACAACA
ATACTGCTAG	CTTTCAGGAT	GATTTTAAA	AAATAGATTG
AAATGTGTTA	TCCTCTCTCT	GAAACGCTTC	CTATAACTCG
AGTTTATAGG	GGAAGAAAAA	GCTATTGTTT	ACAATTATAT
CACCATTAAG	GCAACTGCTA	CACCCTGCTT	TGTATTCTGG
GCTAAGATTG	ATTAAAAACT	AGCTGCTCT	

Fig. 12B